SEQUENCE LISTING

```
Ekwuribe, Nnochiri
       Radhakrishnan, Balasingam
       Price, Christopher
       Anderson, Wesley
       Ansari, Aslam
<120> BLOOD-BRAIN BARRIER THERAPEUTICS
<130> 9233.8DV1
<140> US 09/429,798
<141> 1999-10-29
<150> US 09/134,803
<151> 1998-08-14
<160> 52
<170> PatentIn version 3.2
<210> 1
<211> 6
<212> PRT
     Artificial sequence
<213>
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222> (6)..(6)
<223> Polymer connected to epsilon-amino group
<400>
Tyr Gly Gly Phe Met Lys
1
<210> 2
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
      (1)..(1)
<222>
      Polymer connected to alpha-amino group
<223>
<220>
```

```
<221> MOD RES
<222>
      (6)..(6)
<223> Polymer connected to epsilon-amino group
<400> 2
Tyr Gly Gly Phe Met Lys
                5
<210> 3
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222> (1)..(1)
     Polymer connected to alpha-amino group
<400> 3
Tyr Gly Gly Phe Met Lys
                5
<210> 4
<211> 6
<212>
     PRT
<213> Artificial sequence
<220>
<223>
      Synthetic construct
<220>
<221> MOD RES
<222> (1)..(1)
<223> ACETYLATION
<220>
<221> MOD_RES
      (6)..(6)
<222>
<223>
      AMIDATION
<400> 4
Phe Arg Trp Trp Tyr Lys
1
                5
<210> 5
<211> 6
```

```
<212> PRT
 <213> Artificial sequence
 <220>
 <223> Synthetic construct
 <220>
 <221> MOD_RES
 <222> (1)..(1)
 <223> ACETYLATION
 <220>
 <221> MOD_RES
 <222>. (6)..(6)
 <223> AMIDATION
 <400> 5
Arg Trp Ile Gly Trp Lys
 1
 <210> 6
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
      (6)..(6)
<222>
      AMIDATION
<223>
<220>
<221> MISC_FEATURE
<222> (6)..(6)
<223> Xaa can be any naturally occurring amino acid
<400> 6
Trp Trp Pro Lys His Xaa
<210> 7
<211> 4
<212> PRT
<213> Artificial sequence
<220>
```

<223> Synthetic construct

```
<220>
  <221> MOD_RES
  <222> (4)..(4)
  <223> AMIDATION
  <220>
  <221> MISC_FEATURE
  <222> (4)..(4)
  <223> Xaa is either Lys or Arg
  <400> 7
 Trp Trp Pro Xaa
  1
 <210> 8
 <211> 6
 <212> PRT
 <213> Artificial sequence
 <220>
 <223> Synthetic construct
 <220>
 <221> MOD_RES
 <222> (6)..(6)
 <223> AMIDATION
 <220>
 <221> MISC_FEATURE
 <222> (6)..(6)
 <223>
       Xaa can be any naturally occurring amino acid
 <400> 8
Tyr Pro Phe Gly Phe Xaa
<210> 9
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222> (1)..(5)
<223> Amino acids are in the D-form
<220>
```

```
(6)..(6)
<222>
       n is 0 or 1
<223>
<220>
<221>
      MISC_FEATURE
<222>
      (7)..(7)
<223> Xaa is Gly or the D-form of any naturally occurring amino acid
<220>
<221>
      MOD_RES
<222>
      (7)..(7)
      Amidation
<223>
<400> 9
Ile Met Ser Trp Trp Gly Xaa
<210>
      10
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222>
       (1)..(4)
<223>
      Amino acids are in the D-form
<220>
      MISC_FEATURE
<221>
<222>
      (6)..(6)
      Xaa is Gly or the D-form of any naturally occurring amino acid
<220>
<221> MOD_RES
<222> (6)..(6)
<223> AMIDATION
<400> 10
Ile Met Thr Trp Gly Xaa
1
                5
<210> 11
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
```

```
<220>
      MISC_FEATURE
<221>
       (2)..(2)
<222>
      Xaa is Al, wherein Al is the D-form of Nve or Mle
<223>
<220>
<221>
      MISC FEATURE
      (3)..(3)
<222>
      Xaa is B2, wherein B2 is Gly, Phe, or Trp
<223>
<220>
<221>
      MISC_FEATURE
      (4)..(4)
<222>
      Xaa is C3, wherein C3 is Trp or Nap
<223>
<220>
      MOD RES
<221>
       (4)..(4)
<222>
       AMIDATION
<223>
<400>
      11
Tyr Xaa Xaa Xaa
1
       12
<210>
       3
<211>
      PRT
<212>
       Artificial sequence
<213>
<220>
       Synthetic construct
<223>
<220>
       MOD RES
<221>
<222> (1)..(1)
<223> Tyr has at its N-terminus a Me-x-H-y-N group, wherein x is 0, 1,
       or 2; and y is 0, 1, or 2, with the proviso that x and y is never
       greater than 2
<220>
<221>
       MOD_RES
       (1)..(2)
<222>
<223> The amine between the first Tyr and the second Tyr is methylated,
       wherein z is 0 or 1
<220>
<221> MISC_FEATURE
<222> (3)..(3)
<223> Xaa is Xaa-z, wherein Xaa is Phe, D-Phe or NHBzl, and wherein z
        is 0 or 1
<220>
 <221> MOD_RES
```

```
<222>
      (3)..(3)
<223>
       AMIDATION
<400> 12
Tyr Tyr Xaa
1
<210> 13
<211>
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (4)..(4)
<223> Xaa is D4, wherein D4 is Lys or Arg
<220>
<221>
     MOD_RES
      (5)..(5)
<222>
<223> His is His-z, wherein z is 0 or 1
<220>
<221> MISC_FEATURE
<222>
       (6)..(6)
<223>
       Xaa is Xaa-z, wherein Xaa is any naturally occuring amino acid
       and z is 0 or 1
<220>
<221>
       MOD_RES
<222>
       (6)..(6)
<223> AMIDATION
<400> 13
Trp Trp Pro Xaa His Xaa
<210> 14
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
```

```
Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400> 14
Tyr Xaa Phe Phe
1
<210> 15
<211>
      4
<212> PRT
<213> Artificial sequence
<220>
      Synthetic construct
<223>
<220>
      MISC_FEATURE
<221>
       (2)..(2)
<222>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MOD_RES
      (4)..(4)
<222>
       AMIDATION
<223>
      15
<400>
Tyr Xaa Phe Phe
1
<210>
       16
<211>
       4
<212> PRT
<213> Artificial sequence
<220>
       Synthetic construct
<223>
<220>
<221> MOD_RES
<222> (1)..(1)
       Tyr is Tyr (N-alpha-Me), i.e. N-alpha-methyl tyrosine
<223>
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400> 16
Tyr Xaa Phe Phe
```

```
<210>
      17
<211>
<212>
      PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
      (1)..(1)
<222>
       Tyr is Tyr (N-alpha-Cmp), i.e. N-alpha-cyclopropylmethyltyrosine
<223>
<220>
<221>
      MISC_FEATURE
<222>
      (2)..(2)
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<400> 17
Tyr Xaa Phe Phe
1
<210>
      18
<211>
      4
<212> PRT
<213> Artificial sequence
<220>
<223>
      Synthetic construct
<220>
<221>
      MOD RES
      (1)...(1)
<222>
      Tyr is Tyr (N-alpha-hex), i.e. N-alpha-hexyltyrosine
<223>
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<400> 18
Tyr Xaa Phe Phe
<210> 19
<211> 4
<212> PRT
<213> Artificial sequence
<220>
```

```
<223> Synthetic construct
<220>
<221> MOD_RES
      (1)..(1)
<222>
       Tyr is Tyr (N-alpha-Et2), i.e. N-alpa-diethyltyrosine
<220>
<221>
       MISC_FEATURE
      (2)..(2)
<222>
<223>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400> 19
Tyr Xaa Phe Phe
<210>
       20
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222>
       (1)..(1)
<223>
       Tyr is Dmt, i.e. 2,6-dimethyltyrosine
<220>
<221>
      MISC FEATURE
      (2)..(2)
<222>
<223>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400> 20
Tyr Xaa Phe Phe
<210> 21
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD RES
<222> (1)..(1)
<223> Tyr is Dmt, i.e. 2,6-dimethyltyrosine
```

```
<220>
<221>
       MISC_FEATURE
<222>
       (2)..(2)
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MOD_RES
<222>
       (4)..(4)
<223>
       AMIDATION
<400> 21
Tyr Xaa Phe Phe
1
<210>
      22
<211> 4
<212> PRT
       Artificial sequence
<213>
<220>
<223> Synthetic construct
<220>
<221>
       MOD RES
      (1)..(1)
<222>
       Tyr is H-Tyr(3-F), i.e. 3-fluorotyrosine
<223>
<220>
<221> MISC_FEATURE
<222>
      (2)..(2)
<223>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400>
       22
Tyr Xaa Phe Phe
<210> 23
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD RES
<222>
      (1)..(1)
<223>
      Tyr is H-Tyr(3-Cl), i.e. 3-chlorotyrosine
<220>
```

```
<221>
        MISC_FEATURE
 <222>
       (2)..(2)
 <223>
        Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
 <400> 23
 Tyr Xaa Phe Phe
 1
 <210> 24
 <211> 4
 <212> PRT
 <213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222>
       (1)..(1)
<223>
       Tyr is H-Tyr (3-Br), i.e. 3-bromotyrosine
<220>
<221>
       MISC_FEATURE
<222>
       (2)..(2)
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<400>
      24
Tyr Xaa Phe Phe
<210>
       25
<211>
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD RES
<222> (1)..(1)
       Tyr is Dmt, i.e. 2,6-dimethyltyrosine
<223>
<220>
<221> MISC_FEATURE
<222>
      (2)..(2)
<223> Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
```

```
(2)..(3)
<222>
      nonpetidyl bond
<223>
<400> 25
Tyr Xaa Phe Phe
1
<210>
      26
<211>
      4
<212>
      PRT
<213>
      Artificial sequence
<220>
      Synthetic construct
<223>
<220>
<221> MOD_RES
      (1)..(1)
<222>
       Tyr is Dmt, i.e. 2,6-dimethyltyrosine
<223>
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
<223>
      Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221>
      MOD_RES
       (2)..(3)
<222>
      nonpeptidyl bond
<223>
<220>
<221>
      MOD_RES
<222>
       (4)..(4)
<223> AMIDATION
<400> 26
Tyr Xaa Phe Phe
<210> 27
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
```

```
<223> Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221>
      MOD_RES
<222> (3)..(3)
<223> Phe is -NCH3]Phe, i.e. N-methylphenylalanine
<400> 27
Tyr Xaa Phe Phe
<210> 28
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222>
      (2)..(2)
      Xaa is Tic-psi-[CH2-], i.e.
<223>
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221>
      MOD_RES
      (3)..(3)
<222>
      Phe is -NH] Hfe, i.e. homophenylalanine
<400>
     28
Tyr Xaa Phe Phe
1
<210> 29
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222> (1)..(1)
<223> Tyr is Tyr(NMe), i.e. N-methyltyrosine
<220>
<221> MISC_FEATURE
<222> (2)..(2)
```

```
<223> Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
      MOD_RES
<221>
      (3)..(3)
<222>
     Phe is -NH]Hfe, i.e. homophenylalanine
<223>
<400>
      29
Tyr Xaa Phe Phe
1
<210>
       30
<211>
<212> PRT
      Artificial sequence
<213>
<220>
       Synthetic construct
<223>
<220>
<221>
      MISC FEATURE
<222>
       (2)..(2)
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221>
      MOD_RES
      (3)..(3)
<222>
       Gly is Phg, i.e. phenylglycine
<223>
      30
<400>
Tyr Xaa Gly Phe
1
<210> 31
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<400> 31
Tyr Xaa Trp Phe
```

```
<210> 32
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
      (4)..(4)
<222>
<223> AMIDATION
<400> 32
Tyr Xaa Trp Phe
1
<210> 33
<211> 4
<212> PRT
      Artificial sequence
<213>
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<400> 33
Tyr Xaa His Phe
<210> 34
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
```

```
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
<222> (3)..(3)
<223> Ala is 2-Nal, i.e. 3-(2'-napthyl)alanine
<400> 34
Tyr Xaa Ala Phe
<210> 35
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222>
       (2)..(2)
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MISC_FEATURE
<222> (3)..(3)
       Xaa is Atc, i.e. 2-aminotetralin-2-carboxylic acid
<223>
<400>
      35
Tyr Xaa Xaa Phe
1
<210>
      36
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MOD RES
       (4)...(4)
<222>
```

```
<223> Phe is Phe(pNO2), i.e. 4-nitrophenylalanine
<400> 36
Tyr Xaa Phe Phe
1
<210>
      37
<211>
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
      MISC_FEATURE
<221>
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MOD_RES
<222> (4)..(4)
<223> Phe is Phe(pNO2), i.e. 4-nitrophenylalanine
<400> 37
Tyr Xaa Trp Phe
1
<210>
       38
<211>
       \mathtt{PRT}
<212>
<213> Artificial sequence
<220>
       Synthetic construct
<223>
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
       Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221>
       MOD_RES
      (4)..(4)
<222>
<223> AMIDATION
<400> 38
Tyr Xaa Phe Trp
```

```
<210>
     39
<211>
      7
<212>
     PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
      (7)..(7)
<222>
      AMIDATION
<223>
<400> 39
Tyr Xaa Phe Phe Val Val Gly
                5
<210> 40
<211>
      7
<212>
     PRT
      Artificial sequence
<213>
<220>
<223> Synthetic construct
<220>
<221>
      MISC FEATURE
       (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
<222>
      (7)..(7)
<223>
      AMIDATION
<400> 40
Tyr Xaa Phe Phe Tyr Pro Ser
                5
<210> 41
<211> 7
<212> PRT
<213> Artificial sequence
<220>
```

```
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221>
      MOD_RES
      (7)..(7)
<222>
<223>
      AMIDATION
<400> 41
Tyr Xaa Trp Phe Tyr Pro Ser
<210> 42
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221>
     MOD_RES
      (4)..(4)
<222>
      Phe is Phe(pNO2), i.e. 4-nitrophenylalanine
<223>
<220>
<221> MOD_RES
<222> (7)..(7)
<223> AMIDATION
<400> 42
Tyr Xaa Trp Phe Tyr Pro Ser
<210> 43
<211> 7
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
```

```
<220>
      MISC_FEATURE
<221>
<222>
      (2)..(2)
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
<221> MOD_RES
<222>
      (6)..(6)
<223>
      Nle
<220>
<221>
     MOD RES
      (7)..(7)
<222>
<223>
      AMIDATION
<400> 43
Tyr Xaa Phe Phe Leu Leu Asp
                5
1
<210> 44
<211>
<212>
     PRT
<213> Artificial sequence
<220>
     Synthetic construct
<223>
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<400> 44
Tyr Xaa Phe
1
<210> 45
<211> 3
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
      (2)..(2)
<222>
      Xaa is Tic, i.e. 1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<223>
<220>
```

```
MOD RES
<221>
<222>
       (3)..(3)
<223>
       AMIDATION
<400> 45
Tyr Xaa Phe
1
<210> 46
<211>
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222>
      (2)..(2)
<223>
      Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221>
      MOD_RES
<222>
      (2)..(3)
<223>
      nonpeptidyl bond
<400>
      46
Tyr Xaa Phe
1
<210>
      47
<211> 4
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MISC_FEATURE
<222> (2)..(2)
<223> Xaa is Tic-psi-[CH2-], i.e.
       3-methyl-1,2,3,4-tetrahydroisoquinoline-3-carboxylic acid
<220>
<221> MOD_RES
<222> (2)..(3)
<223> nonpeptidyl bond
<400> 47
```

```
Tyr Xaa Phe Phe
 <210> 48
 <211> 5
 <212> PRT
 <213> Artificial sequence
<220>
<223> Synthetic construct
<400> 48
Tyr Gly Gly Phe Met
<210>
       49
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<400> 49
Tyr Gly Gly Phe Met Lys
<210>
       50
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222> (1)..(1)
<223> NH2 of Tyr is blocked by butyloxycarbonyl group
<400> 50
Tyr Gly Gly Phe Leu Lys
<210> 51
<211> 6
<212> PRT
```

<213> Artificial sequence

```
<220>
 <223> Synthetic construct
<220>
<221> MOD_RES
<222> (1)..(1)
<223> NH2 of Tyr is blocked by butyloxycarbonyl group
<400> 51
Tyr Gly Gly Phe Leu Lys
1
<210> 52
<211> 6
<212> PRT
<213> Artificial sequence
<220>
<223> Synthetic construct
<220>
<221> MOD_RES
<222>
      (1)..(1)
       NH2 of Tyr is blocked by butyloxycarbonyl group
<223>
<220>
<221>
      MOD_RES
<222>
       (6)..(6)
<223> Polymer connected to epsilon-amino group
<400>
       52
Tyr Gly Gly Phe Leu Lys 5
```